





BLAST PubMed Nucleotide Protein Genome Structure PopSet Taxonomy Help

[BLAST with Microbial genome](#)

## BLAST with *Staphylococcus epidermidis* RP62A genome

Total Bases Sequenced: 2616530 bp

Last updated: Jan 9, 2002.

Organism: *Staphylococcus epidermidis* RP62A

Genetic Code: [Translation Table 11](#)

Lineage: Bacteria; Firmicutes; Bacillales; Staphylococcus

Enter your query sequence as GI or FASTA:

```
GVESVRLKILSVIGLLFVL  
IATAACGNNSNSKESSK  
DGVEIKHEEGTTKVPKHPKR
```

Select type of query and database or BLAST-program

Query:  Protein  Database:  Protein  Blast-program:  blastp

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Comments and suggestions to: [\[info@ncbi.nlm.nih.gov\]](mailto:info@ncbi.nlm.nih.gov)



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All: 1 Review: 0

**1:** [Gill SR, Fouts DE, Archer GL, Mongodin EF, Deboy RT, Ravel J, Paulsen IT, Kolonay JF, Brinkac L, Beanan M, Dodson RJ, Daugherty SC, Madupu R, Angiuoli SV, Durkin AS, Haft DH, Vamathevan J, Khouri H, Utterback T, Lee C, Dimitrov G, Jiang L, Qin H, Weidman J, Tran K, Kang K, Hance IR, Nelson KE, Fraser CM.](#)

 Insights on evolution of virulence and resistance from the complete genome analysis of an early methicillin-resistant *Staphylococcus aureus* strain and a biofilm-producing methicillin-resistant *Staphylococcus epidermidis* strain.

J Bacteriol. 2005 Apr;187(7):2426-38.

PMID: 15774886 [PubMed - indexed for MEDLINE]

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